

SEQUENCE LISTING

<110> Nakajima, Toshihiro
Amano, Tetsuya
Yagashita, Naoko

<120> THERAPEUTIC PREPARATION FOR HEMATOPOIETIC DISEASE

<130> BHP-A0301P

<140> US 10/568,033
<141> 2006-02-10

<150> PCT/JP2004/011951
<151> 2004-08-13

<150> US 60/495,001
<151> 2003-08-13

<160> 4

<170> PatentIn Ver. 2.0

<210> 1
<211> 3374
<212> DNA
<213> Homo sapiens

<220>
<221> CDS
<222> (403)..(2256)

<400> 1
gccctttctt atgagcatgc ctgtgttggg ttgacagtga gggtaataat gacttgttgg 60
ttgatttagt atatagggct ctcccttgca aggttaattag gtcctttaaa ttacctgtaa 120
gattttcttg ccacagcatc cattctggtt aggctggta tcttctgagt agtgatagat 180
tggttggtgg tgagggttac aggtgttccc ttctcttact cctggtgttg gctacaatca 240
ggtggcgtct agagcagcat gggacaggtg ggtaaggaaa gtcttctcat tatgcagaag 300
tgatcaactt aaatctctgt cagatctacc tttatgttagc ccggcagtcg cgccggattga 360
gcgggctcgc ggcgctgggt tcctggtctc cgggcccagg ca atg ttc cgc acg 414
Met Phe Arg Thr
1

gca gtg atg atg gcg gcc agc ctg gcg ctg acc ggg gct gtg gtg gct 462
Ala Val Met Met Ala Ala Ser Leu Ala Leu Thr Gly Ala Val Val Ala
5 10 15 20

cac gcc tac tac ctc aaa cac cag ttc tac ccc act gtg gtg tac ctg 510
His Ala Tyr Tyr Leu Lys His Gln Phe Tyr Pro Thr Val Val Tyr Leu
25 30 35

acc aag tcc agc ccc agc atg gca gtc ctg tac atc cag gcc ttt gtc Thr Lys Ser Ser Pro Ser Met Ala Val Leu Tyr Ile Gln Ala Phe Val 40 45 50	558
ctt gtc ttc ctt ctg ggc aag gtg atg ggc aag gtg ttc ttt ggg caa Leu Val Phe Leu Leu Gly Lys Val Met Gly Lys Val Phe Phe Gly Gln 55 60 65	606
ctg agg gca gca gag atg gag cac ctt ctg gaa cgt tcc tgg tac gcc Leu Arg Ala Ala Glu Met Glu His Leu Leu Glu Arg Ser Trp Tyr Ala 70 75 80	654
gtc aca gag act tgt ctg gcc ttc acc gtt ttt cgg gat gac ttc agc Val Thr Glu Thr Cys Leu Ala Phe Thr Val Phe Arg Asp Asp Phe Ser 85 90 95 100	702
ccc cgc ttt gtt gca ctc ttc act ctt ctt ctc ttc ctc aaa tgt ttc Pro Arg Phe Val Ala Leu Phe Thr Leu Leu Phe Leu Lys Cys Phe 105 110 115	750
cac tgg ctg gct gag gac cgt gtg gac ttt atg gaa cgc agc ccc aac His Trp Leu Ala Glu Asp Arg Val Asp Phe Met Glu Arg Ser Pro Asn 120 125 130	798
atc tcc tgg ctc ttt cac tgc cgc att gtc tct ctt atg ttc ctc ctg Ile Ser Trp Leu Phe His Cys Arg Ile Val Ser Leu Met Phe Leu Leu 135 140 145	846
ggc atc ctg gac ttc ctc ttc gtc agc cac gcc tat cac agc atc ctg Gly Ile Leu Asp Phe Leu Phe Val Ser His Ala Tyr His Ser Ile Leu 150 155 160	894
acc cgt ggg gcc tct gtg cag ctg gtg ttt ggc ttt gag tat gcc atc Thr Arg Gly Ala Ser Val Gln Leu Val Phe Gly Phe Glu Tyr Ala Ile 165 170 175 180	942
ctg atg acg atg gtg ctc acc atc ttc atc aag tat gtg ctg cac tcc Leu Met Thr Met Val Leu Thr Ile Phe Ile Lys Tyr Val Leu His Ser 185 190 195	990
gtg gac ctc cag agt gag aac ccc tgg gac aac aag gct gtg tac atg Val Asp Leu Gln Ser Glu Asn Pro Trp Asp Asn Lys Ala Val Tyr Met 200 205 210	1038
ctc tac aca gag ctg ttt aca ggc ttc atc aag gtt ctg ctg tac atg Leu Tyr Thr Glu Leu Phe Thr Gly Phe Ile Lys Val Leu Leu Tyr Met 215 220 225	1086
gcc ttc atg acc atc atg atc aag gtg cac acc ttc cca ctc ttt gcc Ala Phe Met Thr Ile Met Ile Lys Val His Thr Phe Pro Leu Phe Ala 230 235 240	1134
atc cgg ccc atg tac ctg gcc atg aga cag ttc aag aaa gct gtg aca Ile Arg Pro Met Tyr Leu Ala Met Arg Gln Phe Lys Lys Ala Val Thr 245 250 255 260	1182

gat gcc atc atg tct cgc cga gcc atc cgc aac atg aac acc ctg tat		1230
Asp Ala Ile Met Ser Arg Arg Ala Ile Arg Asn Met Asn Thr Leu Tyr		
265	270	275
cca gat gcc acc cca gag gag ctc cag gca atg gac aat gtc tgc atc		1278
Pro Asp Ala Thr Pro Glu Glu Leu Gln Ala Met Asp Asn Val Cys Ile		
280	285	290
atc tgc cga gaa gag atg gtg act ggt gcc aag aga ctg ccc tgc aac		1326
Ile Cys Arg Glu Glu Met Val Thr Gly Ala Lys Arg Leu Pro Cys Asn		
295	300	305
cac att ttc cat acc agc tgc ctg cgc tcc tgg ttc cag cgg cag cag		1374
His Ile Phe His Thr Ser Cys Leu Arg Ser Trp Phe Gln Arg Gln Gln		
310	315	320
acc tgc ccc acc tgc cgt atg gat gtc ctt cgt gca tcg ctg cca gcg		1422
Thr Cys Pro Thr Cys Arg Met Asp Val Leu Arg Ala Ser Leu Pro Ala		
325	330	340
cag tca cca cca ccc ccg gag cct gcg gat cag ggg cca ccc cct gcc		1470
Gln Ser Pro Pro Pro Glu Pro Ala Asp Gln Gly Pro Pro Pro Ala		
345	350	355
ccc cac ccc cca cca ctc ttg cct cag ccc ccc aac ttc ccc cag ggc		1518
Pro His Pro Pro Leu Leu Pro Gln Pro Pro Asn Phe Pro Gln Gly		
360	365	370
ctc ctg cct ctt cct cca ggc atg ttc cca ctg tgg ccc ccc atg		1566
Leu Leu Pro Pro Phe Pro Pro Gly Met Phe Pro Leu Trp Pro Pro Met		
375	380	385
ggc ccc ttt cca cct gtc ccg cct ccc ccc agc tca gga gag gct gtg		1614
Gly Pro Phe Pro Pro Val Pro Pro Pro Ser Ser Gly Glu Ala Val		
390	395	400
gct cct cca tcc acc agt gca gca gcc ctt tct cgg ccc agt gga gca		1662
Ala Pro Pro Ser Thr Ser Ala Ala Leu Ser Arg Pro Ser Gly Ala		
405	410	420
gct aca acc aca gct gct ggc acc agt gct act gct gct tct gcc aca		1710
Ala Thr Thr Ala Ala Gly Thr Ser Ala Thr Ala Ala Ser Ala Thr		
425	430	435
gca tct ggc cca ggc tct ggc tct gcc cca gag gct ggc cct gcc cct		1758
Ala Ser Gly Pro Gly Ser Gly Ser Ala Pro Glu Ala Gly Pro Ala Pro		
440	445	450
ggt ttc ccc ttc cct ccc tgg atg ggt atg ccc ctg cct cca ccc		1806
Gly Phe Pro Phe Pro Pro Trp Met Gly Met Pro Leu Pro Pro Pro		
455	460	465
ttt gcc ttc ccc cca atg cct gtg ccc cct gcg ggc ttt gct ggg ctg		1854
Phe Ala Phe Pro Pro Met Pro Val Pro Pro Ala Gly Phe Ala Gly Leu		
470	475	480

acc cca gag gag cta cga gct ctg gag ggc cat gag cg ^g cag cac ctg Thr Pro Glu Glu Leu Arg Ala Leu Glu Gly His Glu Arg Gln His Leu 485 490 495 500	1902
gag gcc cg ^g ctg cag agc ctg cgt aac atc cac aca ctg ctg gac gcc Glu Ala Arg Leu Gln Ser Leu Arg Asn Ile His Thr Leu Leu Asp Ala 505 510 515	1950
gcc atg ctg cag atc aac cag tac ctc acc gtg ctg gcc tcc ttg ggg Ala Met Leu Gln Ile Asn Gln Tyr Leu Thr Val Leu Ala Ser Leu Gly 520 525 530	1998
ccc ccc cg ^g cct gcc act tca gtc aac tcc act gag ggg act gcc act Pro Pro Arg Pro Ala Thr Ser Val Asn Ser Thr Glu Gly Thr Ala Thr 535 540 545	2046
aca gtt gtt gct gct gcc tcc tcc acc agc atc cct agc tca gag gcc Thr Val Val Ala Ala Ser Ser Thr Ser Ile Pro Ser Ser Glu Ala 550 555 560	2094
acg acc cca acc cca gga gcc tcc cca cca gcc cct gaa atg gaa agg Thr Thr Pro Thr Pro Gly Ala Ser Pro Pro Ala Pro Glu Met Glu Arg 565 570 575 580	2142
cct cca gct cct gag tca gtg ggc aca gag gag atg cct gag gat gga Pro Pro Ala Pro Glu Ser Val Gly Thr Glu Glu Met Pro Glu Asp Gly 585 590 595	2190
gag ccc gat gca gca gag ctc cgc cg ^g cgc cgc ctg cag aag ctg gag Glu Pro Asp Ala Ala Glu Leu Arg Arg Arg Leu Gln Lys Leu Glu 600 605 610	2238
tct cct gtt gcc cac tga cactgccccca gcccgcccc agcctctgct Ser Pro Val Ala His 615	2286
ctttttagca gccctcgctg gaacatgtcc tgccaccaag tgccagctcc ctctctgtct 2346	
gcaccaggga gtagtacccc cagctctgag aaagaggcgg catcccctag gccaagtgg 2406	
aagaggctgg ggttcccatt tgactccagt cccaggcagc catggggatc tcgggtcagt 2466	
tccagccttc ctctccaact cttcagccct gtgttctgct gggccatga aggcagaagg 2526	
tttagcctct gagaagccct cttttcccc caccccttc caggagaagg ggctccccct 2586	
ccaagcccta cttgtatgtg cggagtcaca ctgcagtgcc gaacagtatt agctccgtt 2646	
cccaagtgtg gactccagag gggctggagg caagctatga acttgctcgc tggccaccc 2706	
ctaagactgg tacccatttc cttttcttac cctgatctcc ccagaagcct cttgtggtg 2766	
tggctgtgcc ccctatgccc tgtggcattt ctgcgtctta ctggcaacca cacaactcag 2826	
ggaaaggaat gcctggaggt gggggtgcaag gcggggcagca ctgagggacc ctgccccgcc 2886	

cctcccccca ggccccttc ccctgcagct tctcaagtga gactgacctg tctcacccag 2946
 cagccactgc ccagccgcac tccaggcaag ggccagtgcg cctgctcctg accactgcaa 3006
 tcccagcgcc caaggaaggc cacttctcaa ctggcagaac ttctgaagtt tagaattgga 3066
 attacttcct tactagtgtc ttttggtta aattttgtct tttgaagttg aatgcttaat 3126
 cccggaaag aggaacagga gtgccagact cctggcttt ccagttaga aaaggctcg 3186
 tgccaaggag ggaccacagg agctgggacc tgcctgcccc tgtccttcc cttgggttt 3246
 gtgttacaag agttgttga gacagttca gatgattatt taatttgtaa atattgtaca 3306
 aatttaata gcttaaattt tatatacagc caaataaaaa ctgtcattaa caaaaaaaaaa 3366
 aaaaaaaaaa 3374

<210> 2
 <211> 617
 <212> PRT
 <213> Homo sapiens

<400> 2																		
Met	Phe	Arg	Thr	Ala	Val	Met	Met	Ala	Ala	Ser	Leu	Ala	Leu	Thr	Gly			
1														15				
Ala	Val	Val	Ala	His	Ala	Tyr	Tyr	Leu	Lys	His	Gln	Phe	Tyr	Pro	Thr			
														30				
Val	Val	Val	Ala	Tyr	Leu	Thr	Lys	Ser	Ser	Pro	Ser	Met	Ala	Val	Leu	Tyr	Ile	
														35	40	45		
Gln	Ala	Phe	Val	Leu	Val	Phe	Leu	Leu	Gly	Lys	Val	Met	Gly	Lys	Val			
														50	55	60		
Phe	Phe	Gly	Gln	Leu	Arg	Ala	Ala	Glu	Met	Glu	His	Leu	Leu	Glu	Arg			
														65	70	75	80	
Ser	Trp	Tyr	Ala	Val	Thr	Glu	Thr	Cys	Leu	Ala	Phe	Thr	Val	Phe	Arg			
														85	90	95		
Asp	Asp	Phe	Ser	Pro	Arg	Phe	Val	Ala	Leu	Phe	Thr	Leu	Leu	Phe				
														100	105	110		
Leu	Lys	Cys	Phe	His	Trp	Leu	Ala	Glu	Asp	Arg	Val	Asp	Phe	Met	Glu			
														115	120	125		
Arg	Ser	Pro	Asn	Ile	Ser	Trp	Leu	Phe	His	Cys	Arg	Ile	Val	Ser	Leu			
														130	135	140		
Met	Phe	Leu	Leu	Gly	Ile	Leu	Asp	Phe	Leu	Phe	Val	Ser	His	Ala	Tyr			
														145	150	155	160	

His Ser Ile Leu Thr Arg Gly Ala Ser Val Gln Leu Val Phe Gly Phe
 165 170 175
 Glu Tyr Ala Ile Leu Met Thr Met Val Leu Thr Ile Phe Ile Lys Tyr
 180 185 190
 Val Leu His Ser Val Asp Leu Gln Ser Glu Asn Pro Trp Asp Asn Lys
 195 200 205
 Ala Val Tyr Met Leu Tyr Thr Glu Leu Phe Thr Gly Phe Ile Lys Val
 210 215 220
 Leu Leu Tyr Met Ala Phe Met Thr Ile Met Ile Lys Val His Thr Phe
 225 230 235 240
 Pro Leu Phe Ala Ile Arg Pro Met Tyr Leu Ala Met Arg Gln Phe Lys
 245 250 255
 Lys Ala Val Thr Asp Ala Ile Met Ser Arg Arg Ala Ile Arg Asn Met
 260 265 270
 Asn Thr Leu Tyr Pro Asp Ala Thr Pro Glu Glu Leu Gln Ala Met Asp
 275 280 285
 Asn Val Cys Ile Ile Cys Arg Glu Glu Met Val Thr Gly Ala Lys Arg
 290 295 300
 Leu Pro Cys Asn His Ile Phe His Thr Ser Cys Leu Arg Ser Trp Phe
 305 310 315 320
 Gln Arg Gln Gln Thr Cys Pro Thr Cys Arg Met Asp Val Leu Arg Ala
 325 330 335
 Ser Leu Pro Ala Gln Ser Pro Pro Pro Glu Pro Ala Asp Gln Gly
 340 345 350
 Pro Pro Pro Ala Pro His Pro Pro Pro Leu Leu Pro Gln Pro Pro Asn
 355 360 365
 Phe Pro Gln Gly Leu Leu Pro Pro Phe Pro Pro Gly Met Phe Pro Leu
 370 375 380
 Trp Pro Pro Met Gly Pro Phe Pro Pro Val Pro Pro Pro Pro Ser Ser
 385 390 395 400
 Gly Glu Ala Val Ala Pro Pro Ser Thr Ser Ala Ala Ala Leu Ser Arg
 405 410 415
 Pro Ser Gly Ala Ala Thr Thr Ala Ala Gly Thr Ser Ala Thr Ala
 420 425 430
 Ala Ser Ala Thr Ala Ser Gly Pro Gly Ser Gly Ser Ala Pro Glu Ala
 435 440 445
 Gly Pro Ala Pro Gly Phe Pro Phe Pro Pro Trp Met Gly Met Pro
 450 455 460

Leu Pro Pro Pro Phe Ala Phe Pro Pro Met Pro Val Pro Pro Ala Gly
 465 470 475 480

 Phe Ala Gly Leu Thr Pro Glu Glu Leu Arg Ala Leu Glu Gly His Glu
 485 490 495

 Arg Gln His Leu Glu Ala Arg Leu Gln Ser Leu Arg Asn Ile His Thr
 500 505 510

 Leu Leu Asp Ala Ala Met Leu Gln Ile Asn Gln Tyr Leu Thr Val Leu
 515 520 525

 Ala Ser Leu Gly Pro Pro Arg Pro Ala Thr Ser Val Asn Ser Thr Glu
 530 535 540

 Gly Thr Ala Thr Thr Val Val Ala Ala Ser Ser Thr Ser Ile Pro
 545 550 555 560

 Ser Ser Glu Ala Thr Thr Pro Thr Pro Gly Ala Ser Pro Pro Ala Pro
 565 570 575

 Glu Met Glu Arg Pro Pro Ala Pro Glu Ser Val Gly Thr Glu Glu Met
 580 585 590

 Pro Glu Asp Gly Glu Pro Asp Ala Ala Glu Leu Arg Arg Arg Arg Leu
 595 600 605

 Gln Lys Leu Glu Ser Pro Val Ala His
 610 615

<210> 3
 <211> 3028
 <212> DNA
 <213> Homo sapiens

<220>
 <221> CDS
 <222> (60)..(1910)

<400> 3
 gcagtgcgca ggattgagcg ggctcgccggc gctgggttcc tggtctccgg gccaggca 59

 atg ttc cgc acg gca gtg atg atg gcg gcc agc ctg gcg ctg acc ggg 107
 Met Phe Arg Thr Ala Val Met Met Ala Ala Ser Leu Ala Leu Thr Gly
 1 5 10 15

 gct gtg gtg gct cac gcc tac tac ctc aaa cac cag ttc tac ccc act 155
 Ala Val Val Ala His Ala Tyr Tyr Leu Lys His Gln Phe Tyr Pro Thr
 20 25 30

 gtg gtg tac ctg acc aag tcc agc ccc agc atg gca gtc ctg tac atc 203
 Val Val Tyr Leu Thr Lys Ser Ser Pro Ser Met Ala Val Leu Tyr Ile
 35 40 45

cag gcc ttt gtc ctt gtc ctt ctg ggc aag gtg atg ggc aag gtg		251
Gln Ala Phe Val Leu Val Phe Leu Leu Gly Lys Val Met Gly Lys Val		
50 55 60		
ttc ttt ggg caa ctg agg gca gca gag atg gag cac ctt ctg gaa cgt		299
Phe Phe Gly Gln Leu Arg Ala Ala Glu Met Glu His Leu Leu Glu Arg		
65 70 75 80		
tcc tgg tac gcc gtc aca gag act tgt ctg gcc ttc acc gtt ttt cg		347
Ser Trp Tyr Ala Val Thr Glu Thr Cys Leu Ala Phe Thr Val Phe Arg		
85 90 95		
gat gac ttc agc ccc cgc ttt gtt gca ctc ttc act ctt ctt ctc ttc		395
Asp Asp Phe Ser Pro Arg Phe Val Ala Leu Phe Thr Leu Leu Phe		
100 105 110		
ctc aaa tgt ttc cac tgg ctg gct gag gac cgt gtg gac ttt atg gaa		443
Leu Lys Cys Phe His Trp Leu Ala Glu Asp Arg Val Asp Phe Met Glu		
115 120 125		
cgc agc ccc aac atc tcc tgg ctc ttt cac tgc cgc att gtc tct ctt		491
Arg Ser Pro Asn Ile Ser Trp Leu Phe His Cys Arg Ile Val Ser Leu		
130 135 140		
atg ttc ctc ctg ggc atc ctg gac ttc ctc ttc gtc agc cac gcc tat		539
Met Phe Leu Leu Gly Ile Leu Asp Phe Leu Phe Val Ser His Ala Tyr		
145 150 155 160		
cac agc atc ctg acc cgt ggg gcc tct gtg cag ctg gtg ttt ggc ttt		587
His Ser Ile Leu Thr Arg Gly Ala Ser Val Gln Leu Val Phe Gly Phe		
165 170 175		
gag tat gcc atc ctg atg acg atg gtg ctc acc atc ttc atc aag tat		635
Glu Tyr Ala Ile Leu Met Thr Met Val Leu Thr Ile Phe Ile Lys Tyr		
180 185 190		
gtg ctg cac tcc gtg gac ctc cag agt gag aac ccc tgg gac aac aag		683
Val Leu His Ser Val Asp Leu Gln Ser Glu Asn Pro Trp Asp Asn Lys		
195 200 205		
gct gtg tac atg ctc tac aca gag ctg ttt aca ggc ttc atc aag gtt		731
Ala Val Tyr Met Leu Tyr Thr Glu Leu Phe Thr Gly Phe Ile Lys Val		
210 215 220		
ctg ctg tac atg gcc ttc atg acc atc atg atc aag gtg cac acc ttc		779
Leu Leu Tyr Met Ala Phe Met Thr Ile Met Ile Lys Val His Thr Phe		
225 230 235 240		
cca ctc ttt gcc atc cgg ccc atg tac ctg gcc atg aga cag ttc aag		827
Pro Leu Phe Ala Ile Arg Pro Met Tyr Leu Ala Met Arg Gln Phe Lys		
245 250 255		
aaa gct gtg aca gat gcc atc atg tct cgc cga gcc atc cgc aac atg		875
Lys Ala Val Thr Asp Ala Ile Met Ser Arg Arg Ala Ile Arg Asn Met		
260 265 270		

aac acc ctg tat cca gat gcc acc cca gag gag ctc cag gca atg gac Asn Thr Leu Tyr Pro Asp Ala Thr Pro Glu Glu Leu Gln Ala Met Asp 275 280 285	923
aat gtc tgc atc atc tgc cga gaa gag atg gtg act ggt gcc aag aga Asn Val Cys Ile Ile Cys Arg Glu Glu Met Val Thr Gly Ala Lys Arg 290 295 300	971
ctg ccc tgc aac cac att ttc cat acc agc tgc ctg cgc tcc tgg ttc Leu Pro Cys Asn His Ile Phe His Thr Ser Cys Leu Arg Ser Trp Phe 305 310 315 320	1019
cag cgg cag cag acc tgc ccc acc tgc cgt atg gat gtc ctt cgt gca Gln Arg Gln Gln Thr Cys Pro Thr Cys Arg Met Asp Val Leu Arg Ala 325 330 335	1067
tcg ctg cca gcg cag tca cca cca ccc ccg gag cct gcg gat cag ggg Ser Leu Pro Ala Gln Ser Pro Pro Pro Glu Pro Ala Asp Gln Gly 340 345 350	1115
cca ccc cct gcc ccc cac ccc cca cca ctc ttg cct cag ccc ccc aac Pro Pro Pro Ala Pro His Pro Pro Pro Leu Leu Pro Gln Pro Pro Asn 355 360 365	1163
ttc ccc cag ggc ctc ctg cct cct ttt cct cca ggc atg ttc cca ctg Phe Pro Gln Gly Leu Leu Pro Pro Phe Pro Pro Gly Met Phe Pro Leu 370 375 380	1211
tgg ccc ccc atg ggc ccc ttt cca cct gtc ccg cct ccc ccc agc tca Trp Pro Pro Met Gly Pro Phe Pro Pro Val Pro Pro Pro Pro Ser Ser 385 390 395 400	1259
gga gag gct gtg gct cct cca tcc acc agt gca gcc ctt tct cgg ccc Gly Glu Ala Val Ala Pro Pro Ser Thr Ser Ala Ala Leu Ser Arg Pro 405 410 415	1307
agt gga gca gct aca acc aca gct gct ggc acc agt gct act gct gct Ser Gly Ala Ala Thr Thr Ala Ala Gly Thr Ser Ala Thr Ala Ala 420 425 430	1355
tct gcc aca gca tct ggc cca ggc tct ggc tct gcc cca gag gct ggc Ser Ala Thr Ala Ser Gly Pro Gly Ser Gly Ser Ala Pro Glu Ala Gly 435 440 445	1403
cct gcc cct ggt ttc ccc ttc cct cct ccc tgg atg ggt atg ccc ctg Pro Ala Pro Gly Phe Pro Phe Pro Pro Trp Met Gly Met Pro Leu 450 455 460	1451
cct cca ccc ttt gcc ttc ccc cca atg cct gtg ccc cct gcg ggc ttt Pro Pro Pro Phe Ala Phe Pro Pro Met Pro Val Pro Pro Ala Gly Phe 465 470 475 480	1499
gct ggg ctg acc cca gag gag cta cga gct ctg gag ggc cat gag cgg Ala Gly Leu Thr Pro Glu Glu Leu Arg Ala Leu Glu Gly His Glu Arg 485 490 495	1547

cag cac ctg gag gcc cgg ctg cag agc ctg cgt aac atc cac aca ctg Gln His Leu Glu Ala Arg Leu Gln Ser Leu Arg Asn Ile His Thr Leu 500 505 510	1595
ctg gac gcc gcc atg ctg cag atc aac cag tac ctc acc gtg ctg gcc Leu Asp Ala Ala Met Leu Gln Ile Asn Gln Tyr Leu Thr Val Leu Ala 515 520 525	1643
tcc ttg ggg ccc ccc cgg cct gcc act tca gtc aac tcc act gag ggg Ser Leu Gly Pro Pro Arg Pro Ala Thr Ser Val Asn Ser Thr Glu Gly 530 535 540	1691
act gcc act aca gtt gtt gct gct gcc tcc tcc acc agc atc cct agc Thr Ala Thr Thr Val Val Ala Ala Ser Ser Thr Ser Ile Pro Ser 545 550 555 560	1739
tca gag gcc acg acc cca acc cca gga gcc tcc cca cca gcc cct gaa Ser Glu Ala Thr Thr Pro Thr Pro Gly Ala Ser Pro Pro Ala Pro Glu 565 570 575	1787
atg gaa agg cct cca gct cct gag tca gtg ggc aca gag gag atg cct Met Glu Arg Pro Pro Ala Pro Glu Ser Val Gly Thr Glu Glu Met Pro 580 585 590	1835
gag gat gga gag ccc gat gca gca gag ctc cgc cgg cgc cgc ctg cag Glu Asp Gly Glu Pro Asp Ala Ala Glu Leu Arg Arg Arg Arg Leu Gln 595 600 605	1883
aag ctg gag tct cct gtt gcc cac tga cactgccccca gcccagcccc Lys Leu Glu Ser Pro Val Ala His 610 615	1930
agcctctgct ctttgagca gccctcgctg gaacatgtcc tgccaccaag tgccagctcc ctctctgtct gcaccaggga gtagtacccc cagctctgag aaagaggcgg catccctag gccaagtgga aagaggctgg ggttccatt tgactccagt cccaggcagc catggggatc tcgggtcagt tccagccctc ctctccaact cttcagccct gtgttctgct gggccatga aggcagaagg ttttagcctct gagaagccct cttttcccc caccctttc caggagaagg ggctgcccct ccaagcccta cttgtatgtg cggagtaca ctgcagtgcc gaacagtatt agctcccggtt cccaaagtgtg gactccagag gggctggagg caagctatga acttgctcgc tggcccaccc ctaagactgg tacccatttc cttttcttac cctgatctcc ccagaagcct cttgtggtg ggctgtgcc ccctatgcc tggcattt ctgcgtctta ctggcaacca cacaactcag ggaaaggaat gcctggaggt ggggtgcag gcggcagca ctgagggacc ctgccccgcc cctcccccca ggcccattt ccctgcagct tctcaagtga gactgacctg tctcacccag cagccactgc ccagccgcac tccaggcaag ggccagtgcg cctgctcctg	1990 2050 2110 2170 2230 2290 2350 2410 2470 2530 2590 2650

accactgcaa tcccagcgcc caaggaaggc cacttctcaa ctggcagaac ttctgaagtt 2710
 tagaatttggaa attacttcct tactagtgtc ttttggctta aattttgtct tttgaagttg 2770
 aatgcttaat cccgggaaaag aggaacagga gtgccagact cctggtctt ccagttaga 2830
 aaaggctctg tgccaaggag ggaccacagg agctgggacc tgccctcccc tgcctttcc 2890
 ccttggtttt gtgttacaag agttgttggaa gacagttca gatgattttt taatttgtaa 2950
 atattgtaca aattttaata gcttaaattt tatatacagc caaataaaaaa cttgcattaa 3010
 caaaaaaaaaa aaaaaaaaaa 3028

<210> 4
 <211> 616
 <212> PRT
 <213> Homo sapiens

<400> 4
 Met Phe Arg Thr Ala Val Met Met Ala Ala Ser Leu Ala Leu Thr Gly
 1 5 10 15
 Ala Val Val Ala His Ala Tyr Tyr Leu Lys His Gln Phe Tyr Pro Thr
 20 25 30
 Val Val Tyr Leu Thr Lys Ser Ser Pro Ser Met Ala Val Leu Tyr Ile
 35 40 45
 Gln Ala Phe Val Leu Val Phe Leu Leu Gly Lys Val Met Gly Lys Val
 50 55 60
 Phe Phe Gly Gln Leu Arg Ala Ala Glu Met Glu His Leu Leu Glu Arg
 65 70 75 80
 Ser Trp Tyr Ala Val Thr Glu Thr Cys Leu Ala Phe Thr Val Phe Arg
 85 90 95
 Asp Asp Phe Ser Pro Arg Phe Val Ala Leu Phe Thr Leu Leu Phe
 100 105 110
 Leu Lys Cys Phe His Trp Leu Ala Glu Asp Arg Val Asp Phe Met Glu
 115 120 125
 Arg Ser Pro Asn Ile Ser Trp Leu Phe His Cys Arg Ile Val Ser Leu
 130 135 140
 Met Phe Leu Leu Gly Ile Leu Asp Phe Leu Phe Val Ser His Ala Tyr
 145 150 155 160
 His Ser Ile Leu Thr Arg Gly Ala Ser Val Gln Leu Val Phe Gly Phe
 165 170 175
 Glu Tyr Ala Ile Leu Met Thr Met Val Leu Thr Ile Phe Ile Lys Tyr
 180 185 190

Val Leu His Ser Val Asp Leu Gln Ser Glu Asn Pro Trp Asp Asn Lys
 195 200 205

Ala Val Tyr Met Leu Tyr Thr Glu Leu Phe Thr Gly Phe Ile Lys Val
 210 215 220

Leu Leu Tyr Met Ala Phe Met Thr Ile Met Ile Lys Val His Thr Phe
 225 230 235 240

Pro Leu Phe Ala Ile Arg Pro Met Tyr Leu Ala Met Arg Gln Phe Lys
 245 250 255

Lys Ala Val Thr Asp Ala Ile Met Ser Arg Arg Ala Ile Arg Asn Met
 260 265 270

Asn Thr Leu Tyr Pro Asp Ala Thr Pro Glu Glu Leu Gln Ala Met Asp
 275 280 285

Asn Val Cys Ile Ile Cys Arg Glu Glu Met Val Thr Gly Ala Lys Arg
 290 295 300

Leu Pro Cys Asn His Ile Phe His Thr Ser Cys Leu Arg Ser Trp Phe
 305 310 315 320

Gln Arg Gln Gln Thr Cys Pro Thr Cys Arg Met Asp Val Leu Arg Ala
 325 330 335

Ser Leu Pro Ala Gln Ser Pro Pro Pro Pro Glu Pro Ala Asp Gln Gly
 340 345 350

Pro Pro Pro Ala Pro His Pro Pro Pro Leu Leu Pro Gln Pro Pro Asn
 355 360 365

Phe Pro Gln Gly Leu Leu Pro Pro Phe Pro Pro Gly Met Phe Pro Leu
 370 375 380

Trp Pro Pro Met Gly Pro Phe Pro Pro Val Pro Pro Pro Pro Ser Ser
 385 390 395 400

Gly Glu Ala Val Ala Pro Pro Ser Thr Ser Ala Ala Leu Ser Arg Pro
 405 410 415

Ser Gly Ala Ala Thr Thr Ala Ala Gly Thr Ser Ala Thr Ala Ala
 420 425 430

Ser Ala Thr Ala Ser Gly Pro Gly Ser Gly Ser Ala Pro Glu Ala Gly
 435 440 445

Pro Ala Pro Gly Phe Pro Phe Pro Pro Pro Trp Met Gly Met Pro Leu
 450 455 460

Pro Pro Pro Phe Ala Phe Pro Pro Met Pro Val Pro Pro Ala Gly Phe
 465 470 475 480

Ala Gly Leu Thr Pro Glu Glu Leu Arg Ala Leu Glu Gly His Glu Arg
485 490 495

Gln His Leu Glu Ala Arg Leu Gln Ser Leu Arg Asn Ile His Thr Leu
500 505 510

Leu Asp Ala Ala Met Leu Gln Ile Asn Gln Tyr Leu Thr Val Leu Ala
515 520 525

Ser Leu Gly Pro Pro Arg Pro Ala Thr Ser Val Asn Ser Thr Glu Gly
530 535 540

Thr Ala Thr Thr Val Val Ala Ala Ser Ser Thr Ser Ile Pro Ser
545 550 555 560

Ser Glu Ala Thr Thr Pro Thr Pro Gly Ala Ser Pro Pro Ala Pro Glu
565 570 575

Met Glu Arg Pro Pro Ala Pro Glu Ser Val Gly Thr Glu Glu Met Pro
580 585 590

Glu Asp Gly Glu Pro Asp Ala Ala Glu Leu Arg Arg Arg Arg Leu Gln
595 600 605

Lys Leu Glu Ser Pro Val Ala His
610 615